

REVIEW

The interplay between fascia, skeletal muscle, nerves, adipose tissue, inflammation and mechanical stress in musculo-fascial regeneration

A. Zullo^{1,2}, F.P. Mancini¹, R. Schleip³, S. Wearing⁴, L. Yahia⁵, W. Klingler³

¹ Department of Sciences and Technologies, University of Sannio, Benevento, Italy; ² CEINGE Advanced Biotechnologies, s.c.a r.l, Napoli, Italy; ³ Neurophysiological muscle laboratory, Department of Neurosurgery, Ulm University, Guenzburg, Germany; ⁴ School of Clinical Sciences & Institute of Health and Biomedical Innovation, Queensland University of Technology, Kelvin Grove, Australia; ⁵ Laboratory for Innovation and Analysis of Bio-Performance, École Polytechnique de Montréal, Succursale Centre-Ville Montréal, Quebec, Canada

Muscle and connective tissues are structures that provide stability and movement to organisms belonging to the animal kingdom. Fascia and muscle tissues structurally and functionally integrate to form a musculofascial system with exceptional biomechanical properties, which allow animals to accomplish complex tasks. Mechanical overload through intense exercise or injury, however, may damage these tissues over the course of a lifetime. Although mammalian skeletal muscle and fascia both retain a good regenerative potential in the adulthood, regeneration is very sensitive to alterations in the biochemical and physical environment. In this review, the reciprocal role of fascial tissue and skeletal muscle in their regeneration processes are explored. The involvement of adipose and nervous tissue in the regulation of muscle and fascia regeneration are also revised. It is hypothesised, for the first time, that for effective regeneration of skeletal muscle, both muscle and fascial tissues are necessary, and that nervous and adipose tissues contribute and deeply influence this process.

Key words: Skeletal muscle, Fascia, Musculo-fascial regeneration, Extracellular matrix, Muscle loading, Muscle innervation, Intramuscular adipose tissue

INTRODUCTION

Mammalian skeletal muscle has a complex structure composed by different elements, such as myofibres, connective tissue, blood vessels, nerves, adipose cells and immune cells. Trauma, such as strain injuries, which are very common in sport, damage the muscles. Moreover, chronic degenerative neuromuscular diseases, forced inactivity, and aging can significantly impair muscle structure, mass and functionality¹. Damaged skeletal muscle can regenerate through the coordinated action of different cells and mechanisms, but this process is very complex and unfavorable factors can lead to ineffective muscle repair. Much has been

already understood about skeletal muscle regeneration, but many of its aspects, such as the contribution of fascia, nerves, and adipose tissue still remain poorly investigated.

STRUCTURE OF FASCIAL TISSUE

Fascia is a fibrous connective tissue, which envelopes and infiltrates all the organs and structures of the human body, including muscles, vessels, nerves, and joints². Although the gross anatomical structure of fascia varies throughout the human body, all fascial tissue is composed of cells and an extracellular matrix (ECM).

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■ Correspondence: Alberto Zullo, Department of Sciences and Technologies, University of Sannio, via Port'Arsa 11, 82100 Benevento, Italy - E-mail: albzullo@unisannio.it

FASCIAL CELLS

The main cell types within fascia include fibroblasts, myofibroblasts, fibrocytes, adipocytes, telocytes and migrating white blood cells³⁻⁵. Fibroblasts are mesenchyme-derived cells that produce ECM, release cytokines, and can be transformed, under the certain stimuli, into myofibroblasts⁶. Myofibroblasts express α smooth muscle actin (α SMA) and myosin and have contractile properties⁷. Myofibroblasts, therefore, have the capacity to generate tension in the surrounding ECM, and in addition to secreting cytokines, can also migrate, and respond to chemokines⁸. Fibrocytes are monocyte-derived cells that have features of both macrophages and fibroblasts⁷. Fibrocytes migrate into the injured tissue, where they can differentiate into macrophages or into fibroblasts/myofibroblasts⁷. Adipocytes are connective-tissue cells that store and synthesize fat, and white blood cells belong to the immune system. Telocytes are interstitial cells with mechanotransduction features, which are involved in inter-cellular signaling, survival of satellite cells, angiogenesis, and maintenance of tissue homeostasis. These specific properties of telocytes, make them key elements for tissue regeneration^{9,10}.

FASCIA ASSOCIATED WITH SKELETAL MUSCLE TISSUE

Fascia and muscle tissue strictly cooperate to achieve the higher-order function of the whole human body. The fascia of the muscle tissue can be divided into a superficial fascia, a denser deep fascia, which includes muscular septi (like the lateral septum of the upper arm) and a muscle-specific fascial tissue, which includes three different layers: epimysium, perimysium, and endomysium².

Each of these muscle-specific fascial layers has a different molecular and structural composition¹¹. Endomysium, which envelopes myofibers, contains a matrix layer and a particular structure, the basement membrane. The matrix layer is formed by collagen type I, FN, HA, chondroitin sulfate and dermatan sulfate proteoglycans^{12,13}. The basement membrane is composed mainly of matrix proteins; laminins, type IV and VI collagen, nidogens and the HSPG perlecan¹⁴. These proteins are organized in highly specialized and complex networks that make contacts with integrins and dystroglycan on the surface of myofibers^{13,15,16}. Perimysium, which separates muscle fascicles, is associated with a reticular network composed of collagen fibers, through which it controls the number of mitochondria in the subsarcolemmal space. Epimysium is a thick

collagenous sheath that extends from the tendons and envelops many muscle fascicles. Endomysium is a thin layer of connective tissue that surrounds every single muscle fibre. Epimysium, perimysium, and endomysium are predominantly constituted by collagen type I, II and III, which is synthesized by fibroblasts¹⁷. Cells, via integrins, and extracellular- and basement-membrane proteins, therefore, participate in the formation of a complex 3D network within and surrounding muscle tissue¹⁸.

Fascia acts as a mechanical bridge between different tissues. It transfers mechanical stress, which originates in a specific point of the body, to different areas and tissues. This force transmission allows these different body regions to sense the tensile stimulus and activate proprioceptors and mechanoreceptors¹⁹. Interstitial connective tissue contains and support blood vessels, nerves, and lymphatic tissue, and is responsible for the transfer of mechanical force from the muscle to the skeleton²⁰. Nevertheless, fascia is not just a passive force transmitter. It also regulates mechanical stress by adsorbing, storing and releasing kinetic energy².

The main structural component of fascia, the ECM, reflects the biomechanical environment of muscle tissue, with a structural and a functional role that is based on its constituting proteins: laminin, entactin, fibrinogen, types I and VI collagen, and vitronectin^{21,22}. The ECM of fascia not only provides mechanical support to the muscle, but is also fundamental for the transduction of mechanical tension to muscle cells and of muscle contraction to the surrounding tissues¹¹. At the molecular level, this efficient mechanical communication between different tissues is achieved by means of the ECM, cellular membrane receptors linked to the ECM fibers, and cytoplasmic/cytoskeletal proteins linked to membrane receptors¹¹.

The ECM also governs cell orientation and organization²³ and enhances muscle regeneration by regulating muscle cell proliferation, migration, fragmentation and fusion²⁴. Another important function of the ECM is the regulation of growth factor activity during muscle regeneration, through protein-protein interactions¹⁸. Pathologic changes in the ECM, such as fibrotic ECM, can reduce the regenerative potential of muscle fibers, independently of the myogenic potential itself²⁵. The role of the ECM in maintaining tissue and cellular stability is also highlighted by the phenomenon called anoikis; a programmed cell death process (apoptosis) arising from the interruption of the interaction between ECM and adjacent cells²².

There is a constant balance between the amounts of connective tissue and parenchyma in any tissue²⁶. Healthy skeletal muscle has a high volume ratio of muscle cells to fibroblasts, with fibroblast nuclei comprising

8-15% of all nuclei in the tissue²⁷. For normal tissue turnover, cells also release enzymes that degrade the ECM, such as cathepsins, heparanase, hyaluronidases, and metalloproteinases (MMPs)²². Excessive accumulation of connective tissue, in the form of collagen fibers and myofibroblasts, leads to muscle fibrosis, and diseases, such as Dupuytren Disease²⁸⁻³⁰.

EXTRACELLULAR MATRIX

The ECM fills the extracellular space both within and between each tissue, thus forming the interstitial connective tissue, where it acts as a structural scaffold and functional regulator²². The principal components of the ECM are structural proteins (collagens, laminins, fibronectin, vitronectin, tenascin-c, and elastin), specialized proteins (growth factors, small matricellular proteins and small integrin-binding glycoproteins (SIBLINGS)), glycosaminoglycans and proteoglycans, such as metalloproteinases and cytokines, and water¹⁴. Although the precise composition of the ECM varies between different tissues, collectively the proteins of the ECM confer adhesion, strength, and elasticity to fascial tissues¹⁸. Moreover, they also play an important role in inter-cellular signaling, due to their capacity to store and release growth factors, such as TGFs, and IGFs¹⁴.

STRUCTURAL PROTEINS

Fibronectin (FN) binds to several ECM molecules, including collagen and tenascin, and promotes cell adhesion and migration¹⁴. Also, FN binding to integrins is a crucial step in the fibril assembly process¹⁸. Tenascin C regulates the activity of integrins and syndecan and has a fundamental role in muscle innervation and neural function¹⁴. Elastin forms polymeric elastic fibers that cross-link in an extensible 3D network, which moderate the extensibility of tissues when exposed to low-level force. Collagen is a collective term commonly used to refer to a family of structural proteins of the ECM that provide resistance to tension and link together different ECM proteins. The fibrous components of fascia mainly consist of types I, III, IV, V, VI, XI, XII, XIV, XXI collagen⁴. Vitronectin is an adhesion protein binding cells to the ECM. Laminins, on the other hand, bind to integrins and the dystrophin-glycoprotein complex (DGC)¹⁵.

SPECIALIZED PROTEINS

The ECM is linked to cells by means of adhesion proteins (e.g. integrin receptors, mechanosensitive ion channels and tyrosine kinases), localized on the cell membrane and acting as a bridge between ECM fibers and the intracellular cytoskeleton. Integrin receptors bind to laminins, collagens, fibronectin, tenascin and

link ECM proteins to the intracellular environment, via interaction with actin. This organization enables a fast mechanical communication between the extracellular and the intracellular environments¹⁵. Upon receiving a mechanical input, cells can modify their morphology, 3D orientation, and activity. Therefore, tissue restructuring can be influenced by mechanical stress²⁶.

GLYCOSAMINOGLYCANS AND PROTEOGLYCANS

Heparan sulfate proteoglycans (HSPGs) are components of the ECM and the basement membrane of skeletal muscle. HSPGs form multimeric fibrils and bind to many proteins in the ECM, such as FGF-2, thanks to their negative charges¹⁴. Glycosaminoglycans confer gel properties to the ECM and act as important storage system for growth factors and cytokines²¹. Hyaluronan (HA) is a linear polysaccharide whose main function is to provide lubrication, hydration, and resistance to compression³¹. It is also present as in an aggregated form, which increases the viscosity of ECM and reduces the gliding of connective and muscle tissue layers³².

STRUCTURE OF SKELETAL MUSCLE TISSUE

In mammals, skeletal muscle can account for up to 40% of the total body mass, thus representing the most abundant tissue. It maintains posture and enables voluntary body movements under the control of the cortical areas of the central nervous system.

Skeletal muscle is defined as “striated” muscle because of its band-patterned appearance under microscopic examination³³. Muscle fiber is the smallest functional unit of skeletal muscle. It is a long, multinucleated cylindrical cell composed of a large number of contractile elements, the myofibrils. Skeletal muscle is composed of hundreds to thousands of multinucleated muscle fibers. Many individual muscle fibers are bundled in fascicles by a layer of connective tissue, the endomysium. Fascicles are grouped, in turn, in the muscle by a sheet of connective tissue, the perimysium. Two more connective tissue layers, the epimysium, and the deep fascia, wrap the muscle and separate it from other muscles, respectively³⁴.

Muscle cells establish a connection to the ECM via dystrophin- and integrin-associated complexes^{15 35 36}. These junctions allow the transmission of mechanical traction generated by muscle fibers to the ECM. Another important filament-like protein that contributes to the formation of the bridge between the extracellular matrix and the cytoskeleton of muscle cells is syncoilin³⁷. This protein belongs to the dystrophin-associated protein complex and is localized to the sarcolemma of muscle fibers³⁷. Studies on mice have demonstrated

that Syncoilin is fundamental for the lateral transmission of muscle tension to the extracellular environment³⁷. Disease, forced inactivity, injury, malnutrition, and aging can significantly, and dangerously, deplete the mass and functionality of muscles³⁸. The potential of natural or induced skeletal muscle repair is crucial to re-establish tissue integrity after muscular injury.

REGENERATION OF MUSCLE AND FASCIAL TISSUE

To preserve body integrity and function, both fascia and skeletal muscle tissues act synergistically to aid each other in regeneration. In fact, previous studies have shown that both satellite cells and resident fibroblasts are fundamental for effective regeneration of connective tissue and muscle (Fig. 1). In particular, fibroblasts can enhance satellite cell proliferation and self-renewal³⁹, and skeletal muscle cells participate in the rebuilding of connective tissue¹⁴.

The ECM is a central player in muscle development and regeneration. It is a very dynamic structure that constantly adapts to the environment by continuously switching between degradation and re-assembly. It influences cell proliferation, survival, shape, migration, and differentiation. No wonder its alteration may lead to disease¹⁸.

Metalloproteinases (MMPs) and A Disintegrin and Metalloprotease (ADAMs) are proteases involved in the degradation of ECM proteins. MMPs are produced intracellularly as inactive precursors and are activated by extracellular proteolytic activity. ADAMs, in contrast, are transmembrane proteins located on the membrane of muscle cell¹⁴. The balanced control of MMP activity is achieved by the secretion of tissue inhibitors of MMPs (TIMPs) that allow MMP action to be finely tuned¹⁴. Proteolytic degradation of the ECM is also a crucial step for the movement of endothelial cells during regenerative neovascularization⁴⁰. ECM degradation, therefore, not only destroys the extracellular protein network but also generates bioactive peptides and releases growth factors trapped within the ECM¹⁸.

Given the strong interconnection between ECM and skeletal muscle, it is not surprising that skeletal muscle influences the remodeling of the ECM. In fact, during skeletal muscle regeneration extracellular proteases (MMP-2, MMP-7, MMP-9, MMP-24, MMP-25, plasmin and the plasminogen activation system), secreted by muscle cells, inflammatory cells and other cells, participate in the degradation of the ECM⁴¹. MMPs can both degrade ECM proteins, such as collagen fibers, elastin, proteoglycans, and fibronectin as well as proteolytically activate other proteins, such as the MMPs

themselves⁴². For example, plasmin can degrade fibronectin and laminin and also activate MMPs and several latent growth factors⁴³. Moreover, there is some evidence that activated satellite cell influence the remodeling of ECM^{14,25}.

The formation of new muscle mass is a process involving muscle cells and non-muscle cells with myogenic potential, such as adipose-tissue derived stem cells, mesoangioblasts, pericytes, side-population cells, Ac133 + cells, stem and/or precursor cells from muscle endothelium, and synovium⁴⁴. Nevertheless, skeletal muscle regeneration relies mainly on the activity of satellite cells, i.e. muscular stem cells (MuSCs) with asymmetric division and self-renewal capabilities⁴⁵. These cells are small, have a single nucleus, and are located between the myofiber sarcolemma and the basal lamina, and are, therefore, in close contact with the ECM. The paired box 7 (Pax7) transcription factor and tetraspanin KAI/CD82 are the molecular markers that identify satellite cells^{46,47}. Pax7+ satellite cells are capable of asymmetric division, thus contributing to the regeneration of muscle fibers and, at the same time, maintenance of the stem cell pool⁴⁸. Furthermore, MuSCs express different molecular markers in the quiescent state and the proliferating/differentiating state⁴⁶. Even though satellite cells constitute the vast majority of MuSCs, it has been observed that non-satellite cells like mesenchymal stem cells may also acquire the skeletal muscle features under appropriate conditions⁴⁹. Upon skeletal muscle injury, satellite cells are activated and start proliferating to execute the dual program of self-renewing and differentiation into new muscular fibers⁵⁰. The critical balance between post-replicative differentiation and stemness is regulated by the presence of myogenic and self-renewing subpopulations of satellite cells⁵¹. On the other hand, the satellite niche, which is the MuSCs microenvironment, provides specific signals driving satellite cells toward either differentiation or stemness fate⁴⁶. These signals include cell-to-matrix and cell-to-cell interactions, autocrine and paracrine stimuli, as well as other biochemical factors.

Satellite cells live in a reversible 'G0' or quiescence state that is crucial for the maintenance of the pool of myogenic precursors. This state allows satellite cells to re-enter the cell cycle in response to specific stimuli, like muscular injury. Cell cycle entry and exit are controlled by the coordinated expression of an array of regulatory genes and factors. While genes inhibiting the cell cycle are highly expressed in quiescent satellite cells, those promoting the cell cycle are activated after muscular injury⁵²⁻⁵⁵. During muscle regeneration, proliferating satellite cells either exit the cell cycle to return to a quiescent state and restore the MuSC pool or, alternatively, differentiate into myoblasts and fuse with pre-existing

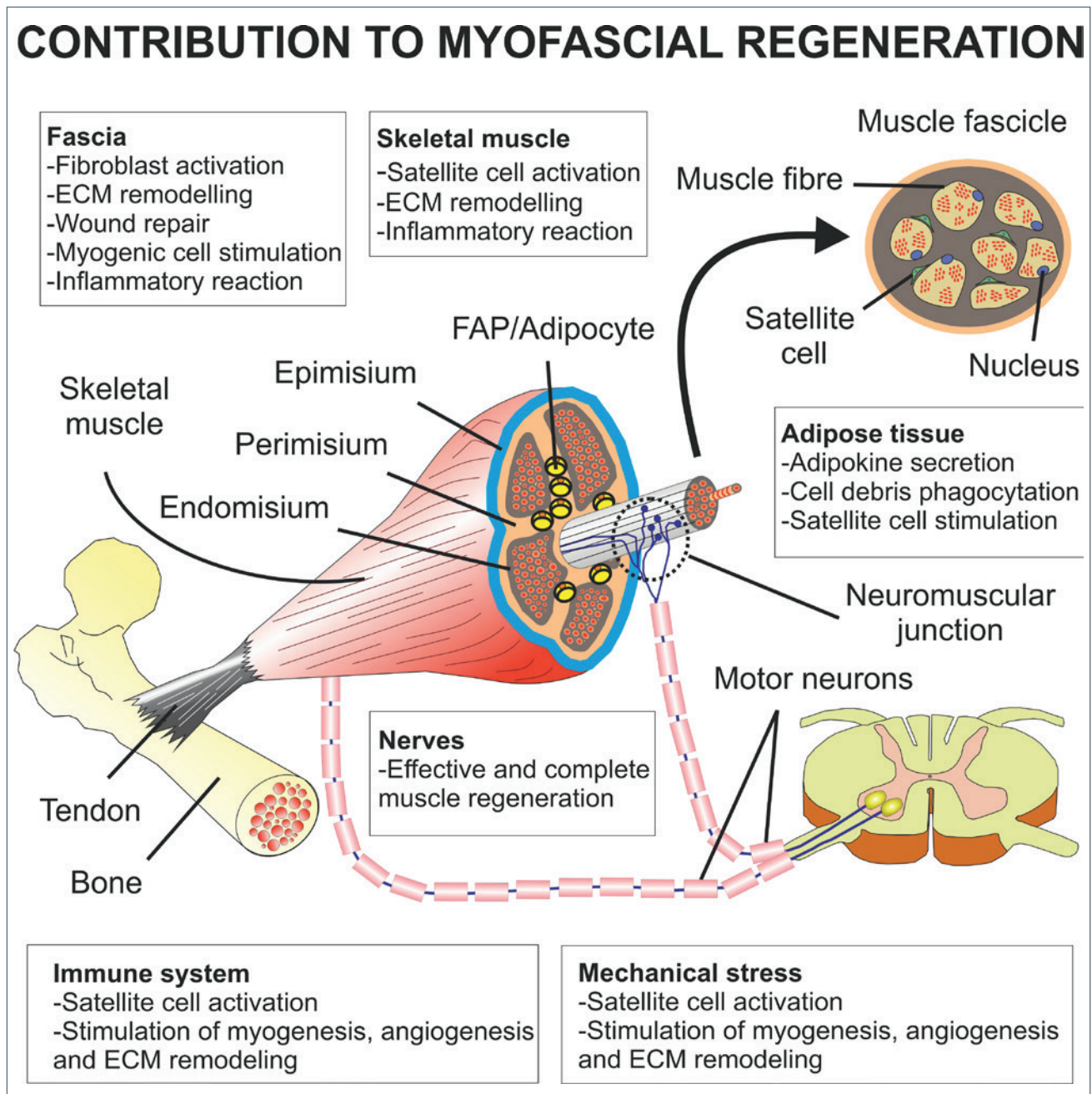


Figure 1. Scheme showing the contribution of skeletal muscle, fascia, nerves, intramuscular adipose tissue, immune system, and mechanical stress to myofascial regeneration.

muscle fibers to form new myofibers^{56,57}. Collagen type VI, present in the ECM, and fibronectin, produced by activated satellite cells, induces the self-renewal of the satellite pool^{58,59}. The post-mitotic cellular contacts within the niche also play a role in the determination of the satellite cell fate. For example, daughter cells in contact with the basement membrane are directed to the satellite cell self-renewal pathway, while daughter

cells interacting with myofibers differentiate into myofibers⁶⁰.

ECM components also enhance muscle regeneration by regulating muscle cell proliferation, migration, fragmentation, and fusion²⁴. Aberrant ECM remodeling after muscle injury, therefore, leads to ineffective muscle regeneration⁶¹. In the initial phase of skeletal muscle regeneration, both fibroblasts and satellite cells actively

proliferate, promoting each other's growth through a reciprocal positive feedback ³⁹.

Cellular activities, as well as temporal and spatial production of ECM proteins, vary according to the different phases of the regeneration process ¹⁴. Following tissue injury, fibroblasts migrate to the site of damage, proliferate and generate an extracellular network of disorganized collagen III fibers, most of which are subsequently transformed into collagen I fibers. Moreover, fibroblasts also secrete growth factors for the myogenic cells and differentiate into myofibroblasts, which contribute to the formation of new muscle fibers ³⁹. At this stage, the ECM collagen type I/III, which confer tensile strength, and laminin, tenascin-C, and fibronectin, which confer elastic properties, are produced, and TGF- β 1 is one important driver of such processes ⁶². After this phase, a wound contracture process shortens the edges of the damaged area ⁶³. The shortening of these margins takes place because of the crosslinking and shortening of the collagen fibers. As result, a tight scar forms, thus increasing the mechanical resistance, and the functional performances of the damaged tissues. Afterward, collagen fibers aggregate and orientate to give better functional responses to mechanical stress, and to contractile activity ⁶³. This fibrotic response is beneficial in the initial phase of tissue regeneration because it preserves the tissue integrity. On the other hand, fibrotic elements should be removed, little by little, from the injured site, to replace the fibrotic tissue with functional skeletal muscle tissue ⁶².

During the regenerative phase, the regrown myofibers establish new and stronger lateral contacts (increasing the numbers of integrins on the adhesion sites) with the surrounding ECM, to allow some muscle activity and reduce the risk of a re-rupture ¹⁴. The role of ECM in regenerative processes is so important that it has also been demonstrated in nonmammalian animal models, such as axolotl and *Octopus Vulgaris* ^{64 65}.

ADIPOSE TISSUE

There is a close physical and biochemical link between connective tissue, muscle tissue, and adipose tissue. Therefore, they profoundly influence each other. In particular, skeletal muscle activity and regeneration are strongly affected by adipose tissue. Adipocytes are present in muscle tissue as "intramuscular fat", and secrete adipokines, proteic hormones that also regulate skeletal muscle development and function (Fig. 1).

Different cell types, including mesenchymal progenitors and fibro/adipogenic progenitors (FAPs), have been shown to participate in the development of fat cells within the muscle tissue, however, the potential of

satellite cells to generate intramuscular adipocytes has only been suggested ⁶⁶. FAPs are multipotent mesenchymal progenitor cells that are present in many tissues, such as skeletal muscle. These cells are fundamental in the process of muscle regeneration. In fact, after muscle injury, they transiently proliferate, phagocytize necrotic debris, and stimulate satellite cell expansion. Importantly, after this initial phase, the number of FAP must rapidly decline to allow efficient muscle regeneration ^{67 68}. Under disuse or chronic muscle diseases, such as dystrophic conditions, FAPs proliferate, differentiate into adipocytes and ECM-producing fibroblasts and persist in the tissue; thereby resulting in the accumulation of intra-muscular adipose tissue (IMAT) and fibrotic tissue ⁶⁹⁻⁷¹. In particular, FAPs can be induced to differentiate into ECM-producing cells by macrophages ⁷². It has been shown in mouse models, that unloading inhibits IMAT accumulation in regenerating muscle ⁷³. The accumulation of fat tissue in the muscle inhibits its regenerative potential because it suppresses the proliferative capacity of satellite cells and may also inhibit the activity of macrophages ^{73 74}. In fact, obese mice have a very low skeletal muscle regenerative potential ⁷⁴. Fatty infiltration in the muscle tissue has also been proposed to arise from the differentiation of FAP into adipocytes when muscle regeneration is impaired ⁶⁷. The crosstalk among MuSCs, FAPs, and immune cells determines the fate of the muscle tissue; either toward regeneration or fibro-adipogenic degeneration ⁷⁵.

In vitro, adipose stem cells can influence the activity of myofibroblasts. In particular, this cell-cell interaction leads to inhibition of actin expression and contractility in smooth muscle ⁷⁶. In the rat, it has been demonstrated that adipose tissue-derived regenerative cells (ADRCs) can stimulate the repair of injured skeletal muscle tissue, through their paracrine activity ⁷⁷. In particular, these cells promote angiogenesis and myogenesis and prevent fibrosis. Moreover, adipose-derived stem cells (ASCs) have been shown *in vitro* to differentiate into skeletal muscle cells ⁷⁸. Experimental evidence has also demonstrated that human ADSC can be transdifferentiated into fibroblast-like cells *in vitro* ⁷⁹. These observations underline the close interconnection between adipose and connective tissue.

The interaction among different tissues also relies on soluble factors that are released from one tissue and target other tissues. Adipose tissue, once believed to be merely an energy depot, secretes an extensive repertoire of molecules, collectively called adipokines, which can affect surrounding cells or distant cells, thus acting in a paracrine or endocrine fashion. Skeletal muscle expresses receptors for several adipokines, including adiponectin and leptin ⁴⁴. Globular adiponectin (gAD) is also involved in the proliferation and myogenic

differentiation of satellite cells, myoblasts, and mesoangioblasts⁴⁴. gAd is also fundamental for the migration of satellite cells and macrophages towards the muscle tissue. Moreover, gAD causes the degradation of the extracellular matrix through the induction of MMP-2 secretion by satellite muscle cells⁴⁴. Adiponectin is also secreted in an autocrine fashion by skeletal muscle tissue, and the inflamed environment stimulates the muscle production of gAd⁴⁴. In this context, the importance of adiponectin is further underlined by the evidence that decreased plasma adiponectin is associated with reduced regenerative potential of skeletal muscle⁴⁴.

Studies on mice have also demonstrated that leptin affects muscle tissue by increasing skeletal muscle mass and altering its miRNA expression⁸⁰. Similarly, Il-6, an adipo-myokine that is secreted both by adipose and muscle tissue has been shown to stimulate satellite cell activation and muscle repair at the muscle level⁸¹. It has also been recently discovered that a novel adipokine, CTRP3, is involved in the proliferation and differentiation of myoblasts. CTRP3 is expressed in muscle cells and acts in an autocrine/paracrine fashion⁸². In addition, other adipokines have also been shown to regulate myogenesis during muscle regeneration⁸².

HUMORAL FACTORS

Soluble factors participate in the signaling process between the immune system, fascial tissue, and muscle tissue. HGF, FGFs, TGF β s, MSTN, IGF, TNF α , PDGF, IL-6, LIF, VEGF, EGF, KGF, are secreted by different cell types, including fibroblasts, myofibroblasts, muscle cells and immune cells²¹. Other factors, such as TGF- β 1, are trapped in the ECM via specific binding proteins (decorin and biglycan) and can be released by mechanical triggers^{18 21 83 84}.

In adult skeletal muscle, TGF- β 1 hinders muscle regeneration and promotes muscle atrophy and fibrosis. In particular, TGF- β 1 inhibits satellite cell proliferation, myofiber fusion, the activation of muscle-specific genes and, after injury, induces myoblast to myofibroblast conversion⁸⁵⁻⁸⁷. Furthermore, neutralization of TGF- β 1 reduces scar tissue formation and improves muscle tissue healing⁶².

IMMUNE CELLS AND INFLAMMATION

The regenerative process in skeletal muscle also requires the participation of non-satellite cells, such as perivascular and immune cells (Fig. 1)⁸⁸. Inflammatory cytokines (IFN-g and TNF), released at the site of injury, induce the polarization of infiltrated monocytes to M1

macrophages (classically proinflammatory). These cells, in turn, release other inflammatory cytokines and IGF-1, which activate muscle satellite cells^{89 90}. Later, changes in the biochemical environment (cytokines) at the site of injury, induce the polarization of M1 macrophages to M2 anti-inflammatory phenotype. M2 macrophages inhibit inflammation by secreting IL-4, IL-10, TGF- β 1, and support myogenesis by secreting IGF-1⁹⁰. M2 macrophages also induce angiogenesis and promote extracellular matrix remodeling⁹⁰⁻⁹². The role of monocytes in the regeneration process of skeletal muscle has been demonstrated in mouse models, where a reduction of monocytes in the bloodstream or skeletal muscle has been shown to impair the regenerative potential of muscle^{93 94}. In a mouse model of muscular dystrophy, suppression of proinflammatory macrophage activity by IL-10 reduces the severity of the disease⁸⁹. This switch from M1 inflammatory to M2 anti-inflammatory and pro-regenerative macrophages is critical for an effective regeneration of the skeletal muscle⁹⁴⁻⁹⁶. In the early phases following muscle injury, inflammation favors muscle regeneration by inhibiting satellite cell death and promoting myoblast proliferation⁴⁴. In the later phases, in contrast, Treg cells enhance the regenerative potential of skeletal muscle by protecting injured tissue from inflammatory damage, promoting satellite cells differentiation (by amphiregulin), and inhibiting fibrosis^{97 98}. During chronic inflammation, such as those occurring in muscular dystrophies, the ECM accumulates and replaces muscle tissue with fibrotic tissue⁶⁷. During muscle regeneration, anti-inflammatory macrophages induce fusion, growth, and differentiation of muscle cells by TGF- β , low levels of TNF- α and IGF-1^{99 100}.

The dense network of the ECM acts as a physical barrier against the movements of cells. Therefore it must be weakened by fragmentation to allow cells to penetrate it. The binding of integrins with laminins and type IV collagen on the basal lamina prevents neutrophils from reaching the skeletal muscle¹⁴. In the area of damage, different cells, such as fibroblasts, macrophages, endothelial cells release several proteases that fragment the ECM¹⁴. The ECM also has a positive role for cell invasion into the injury site. Fibronectin, type IV collagen, laminin, entactin and proteolytic fragments of fibronectin and laminin, all act as guides and chemoattractants for immune cells and myoblasts^{14 101}.

Interestingly, in aged organisms, the mass and strength of skeletal muscle are reduced, and satellite cells have impaired regenerative potential. Although the myogenic commitment of satellite cells is not affected by aging, their capacity to self-renew is reduced¹⁰². Emerging evidence suggests that changes occurring in the satellite cell niche contribute to the reduced regenerative potential of aged muscle.

MECHANICAL STRESS

Mechanical stress has a reciprocal influence on muscle and ECM. The ECM influences muscle cell alignment and activity, and muscle activity influences collagen fiber alignment²⁶. It is well known that physical exercise is healthy for the organism, but only recently has it been appreciated that mechanical loading of skeletal muscle post-injury also triggers many beneficial responses that enhance skeletal muscle regeneration and reduce fibrosis (Fig. 1)¹⁰³⁻¹⁰⁷. Skeletal muscle loading can directly increase skeletal muscle regeneration by stimulating the activation of SCs, attracting exogenous myogenic stem cells and stimulating angiogenesis^{106 107}. Moreover, mechanical loading of newly formed muscle fibers is crucial for maturation of the tissue and of the myotendinous junctions. Interestingly, mechanical stimulation, in the form of stretch activation, massage therapy, and physical manipulation can improve healing of muscle tissue after damage, exerting anti-inflammatory and anti-fibrotic effects¹⁰⁸⁻¹¹⁰.

The formation of scar tissue, secondary to an injury or pathogenic process, modifies the biomechanical properties of the connective tissue. It also leads to a reduction in gliding between adjacent tissues⁶³. Several studies have shown that physical exercise in Duchenne muscle dystrophy and skeletal muscle-injured mice can prevent fibrosis and enhance muscle performance¹¹¹⁻¹¹⁴. Parenthetically, removal of stress, via mechanical unloading, has been shown to reduce muscle regeneration after injury¹¹⁵⁻¹¹⁷. After this initial resting phase, however, the absence of mechanical loading reduces the extent of muscle regeneration⁷³. As demonstrated in ovariectomized mice, mechanical stimulation of muscle fibers can inhibit fat accumulation in skeletal muscle tissue, a side effect of ovariectomy⁶⁶. Mechanical stimulation has also been proved to prevent satellite cells differentiation toward an adipogenic fate⁶⁶.

The contractile activity of muscle cells stimulates the production of myokines, such as IL-6¹⁰⁷. Transient and short-term exercise-induced production of IL-6 is beneficial for muscle metabolism, however, long-term IL-6 production is detrimental and is associated with muscle atrophy¹¹⁸.

Myokines are proteins secreted from skeletal muscle cells that exert autocrine and endocrine effects and allow communication with adipose tissue, liver, and pancreatic cells¹⁰⁷.

Mechanical stress is known to influence the structure the ECM. Deposition and alignment of the collagen fibers secreted into the extracellular space (to form the ECM matrix) is driven by the position and the orientation of the connective fibroblast-like cells²⁶. At molecular

level, the alignment of intracellular actin fibers in collagen-producing cells guides the parallel deposition of extracellular collagen²⁶. Therefore, the stress produced by cytoskeletal proteins of cells can influence the structure of ECM¹⁸.

Mechanical load also influences fiber alignment by acting on ECM protein synthesis and degradation²⁶. Strain-stabilization, a mechanism relating mechanical stress and collagen fiber orientation, involves mechanical activation of enzymatic degradation of the collagen fibers²⁶. ECM fiber orientation, in turn, affects cell activity, differentiation, and migration^{119 120}. Mechanical stress, in the form of traction, also stimulates the regeneration of deep fascia in animals subjected to leg lengthening^{28 121}. Shear stress applied to the tissue can also modify the alignment of collagen fibers²⁶.

The sensitivity of tissues to mechanical stress is dependent on the presence of TGF- β 1 in traction-sensitive protein complexes that can activate/deactivate TGF- β 1 in response to contractile force changes⁸³. These mechanical stresses can originate in the extracellular environment (injury) or by cell contraction (myofibroblasts or muscle cells)⁸³. The mechanical communication between extracellular and intracellular environments is due to specific cell-ECM and cell-cell interactions mediated by protein complexes¹²². Among these, integrins function as mechanotransducers that can transmit tension in the ECM to the cell and activate cellular responses¹⁸. Cells can sense changes in ECM stiffness and mechanical stress by cytoskeletal actomyosin contraction via integrin¹²². Proteins of the ECM interact with cells by integrins, assembled with other proteins in supramolecular adhesion complexes. The extracellular mechanical stress is transduced into intracellular events, by specific integrin-actin cytoskeleton interactions. Mechanical stimuli can alter the membrane assembly of these adhesion complexes and activate specific signaling processes, such as RhoA-induced actin stress fiber formation¹²³⁻¹²⁷.

The input signals for the remodeling of the ECM are transmitted by ECM receptors, such as integrins, laminin receptors, syndecans, and proteases, including MMPs, serine protease (e.g. plasmin, plasminogen activator and uPAR) and cysteine protease (e.g. cathepsins) families and by structural tension provided by cells and extracellular proteins¹⁸.

In the integer tissue, there is always a tensional homeostasis between cell traction and ECM stiffness¹²². There are distinct intracellular protein complexes, such as SPARC-ILK and Rho-A, that can also provide the tension required for FN assembly at the cell surface¹⁸. Moreover, FN can also induce cell contraction¹⁸.

NERVOUS TISSUE

Reinnervation of injured skeletal muscle is crucial for effective and complete regeneration of the tissue (Fig. 1)¹⁴. In particular, the formation of neuromuscular junctions (NMJs) drives gene expression in the regenerating myotubes^{128 129}. The complex ECM network acts as tissue stabilizer and guides the formation of new NMJs¹³⁰. Therefore, ECM is essential for reinnervation of injured skeletal muscle¹⁸. Somewhat surprisingly, however, the presence of nerves does not appear to be critical in the initial phases of muscle regeneration¹⁴. After muscle injury, the regeneration process restores muscle tissue, ECM, nerves, and their connections. Due to specific proteins, such as agrin and s-laminin and components of the basal lamina (BL), skeletal muscle retains the memory of the original NMJ sites and can guide the formation and differentiation of new NMJs at same location as in pre-injured muscle^{14 131}. Integrity of the BL surrounding muscle fibers and motor axons, however, is crucial for successful re-innervation and regeneration of muscle tissue¹⁴. The BL surrounding the nerve can act as a scaffold and a guide axon regeneration toward the original NMJ site. When the continuity of the BL is disrupted, re-innervation of the regenerating muscle is less efficient¹⁴. Nevertheless, even in the situation of BL disruption, NMJs can readily form at the original NMJ sites when contacted by a growing axon¹⁴.

CONCLUSIONS

Regeneration is the evolutionary key to preserve tissue function during aging and after injury. In humans, different tissues and organs have different regenerative potentials. However, tissues with substantial regenerative capacity can fail to regenerate because of disease or antagonistic physical and biochemical factors. Skeletal muscle and fascial tissues have a remarkable ability to regenerate. Nevertheless, their regeneration is strongly dependent on their close interaction and the presence of nervous and adipose tissues. Moreover, the regeneration process can be easily hindered in favor of non-functional fibrotic tissue accumulation. This, in turn, results in loss of the original mechanical and biological function of the injured tissue, increasing the susceptibility to re-injury, and the development of chronic pain^{110 132}. ECM and fibroblasts activity play a fundamental role in the reaction of skeletal muscle to injury¹³³. Regenerative medicine aims to improve recovery and regeneration of damaged tissues. Over the last decade, cell therapy and bio-engineered synthetic scaffolds have been employed to enhance tissue

repair^{18 22 133-136}. These approaches aim to replace damaged tissue with new functional tissue by supplying ECM-like structures with active cells¹⁸.

Although the results of regenerative medicine trials are still not sufficiently encouraging, a deeper comprehension of the complex mechanisms underlying tissue regeneration can significantly improve the outcome of future clinical studies.

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